

FIG.1C

FIG.1D

c12 c21 c23 c40 actin

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FIG. 2A

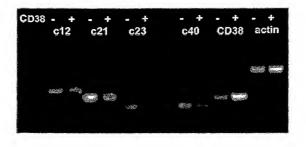


FIG.2B

			1	2	3	4	5	6	7	8
		Α	Heart	Aorta	Skeletai Muscle	Colon	Bladder	Uterus	Prostate	Stomach
		В	Testis	Ovary	Pancreas	Pituitary	Adrenal	Thyroid		Mammary
FIG.3A		С	Kidney	Liver	Small Intestines	Spleen	Thymus	WBC	Lymph Node	Bone Marrow
		D	Appendix	Lung	Trachea	Placenta		Brain	Medulla	Spinal Cord
		Ε	Fetal Brain	Fetal Heart	Fetal Kidney	Fetal Liver	Fetal Spieen	Fetal Thymus	Fetal Lung	
		F	Yeast total RNA 100ng	Yessi s-RNA 100ng	E pad rRNA 100 mg	E coff CNA 100 ng	Poly (A)	Hunton C _{et} 1 DNA 100 ng	Human DNA 100 ng	Human DNA 500 ng
	1		1	2	3	4	100ms 5	6	7	8
		Α							-1	
		В				*	•	•	*	
FIG.3B		С		•		*			*	*
		D								
		Ε						e		
		F	-							
			1	2	3	4	5	6	7	8
		Α	- 1	*		*	•	*	*	٠
		В					•	*	٠	•
FIG.3C	Ì	С	-		٠	•	*		٠	
	1	D			•	*				
		Ε		•	♣,	٠	٠		*	
	Ī	F								
			1	2	3	4	5	6	7	8
		Α	*		,	•			*	*
	I	В			•	*	*		*	
FIG.3D	İ	С	•	*	*		٠			*.
		D	-	*		•		. •		
		Е			*	•				
		F								

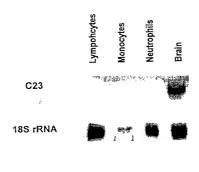


FIG.4A

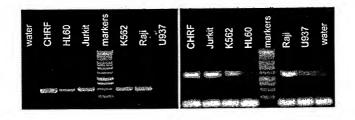


FIG.4B

FIG.4C

```
cogggagggg ggagcggcgt tggaggccac cgtttccagc atcaacaaca
   1
      geaacttgtg attggeggtg aceggatatt cagttgcaca tececacate
  51
      aatgcactgc caatgggtta tatcctgtgt tgtgacctca tggtttaagt
 101
 151
      gggaataaag ATGAGTATAA GCAGTGATGA GGTCAACTTC TTGGTATATA
      GATACTIGCA AGAGICAGGA TITICICATI CAGCATITAC CTITGGIATA
 201
      AAAAGCCATA TCAGTCAGTC CAATATAAAT GGTGCCCTCG TCCCACCCGC
 251
      TGCATTGATT TCTATCATCC AGAAAGGTCT ACAGTATGTA GAAGCAGAAG
 301
 351
      TTAGTATTAA TGAGGATGGT ACCTTGTTTG ATGGTCGACC AATAGAGTCT
 401
      CTGTCCCTGA TAGATGCCGT AATGCCTGAT GTAGTACAAA CAAGACAACA
 451
      AGCTTATAGA GATAAGCTTG CACAGCAACA GGCAGCAGCT GCTGCAGCTG
 501
      CCCCACCTCC ACCCACCCAA CAACGATCTC CAAAAAATGC AGAAAACACA
 551
      CCAAATGGGG AGGAGAATGG AGCACATACT ATAGCAAATA ATCATACTGA
      TATGATGGAA GTGGATGGGG ATGTTGAAAT CCCTCCTAAT AAAGCTGTTG
 601
 651
      TGTTGCGGGG CCATGAATCT GAAGTTTTTA TCTGTGCCTG GAACCCTGTT
      AGRICATORO TAGGATUACO GUETOGACAO TORACAGUA GAAURUGGAA
 701
 751
      TCTTAGTGAG AACAGCACCA GTGGCTCTAC ACAGTTAGTA CTTAGACATT
      CTATACCACA ACCAGGGCAA CATCTTCCGA GCAACAAGGA TGTCACATCT
 801
      CTAGATTGGA ATAGTGAAGG TACACTTCTA GCAACTGGTT CCTATGATGG
 851
 901
      GTTTGCCAGA ATATGGACTA AAGATGGTAA CCTTGCTAGC ACCTTAGGGC
      AGCATAAAGG CCCTATATTT GCATTAAAAT GGAATAAGAA AGGAAATTTC
 951
     ATTEMATOR CITEGRATIACA CANCACTACA ATTAUTIGGG ACCCACATAC
1001
      TGGTGAAGCC AAGCAACAGT TTCCTTTTCA TTCAGCACCA GCATTGGATG
1051
1101
      TTGATTGGCA GAGCAACAAC ACCTTTGCTT CTTGTAGTAC AGATATGTGC
      ATTCATGTCT GTAAATTAGG ACAAGACAGA CCTATTAAAA CATTCCAAGG
1151
1201
     ACATACGAAT GAAGTAAATG CTATCAAATG GGACCCAACT GGCAATCTCT
      TOGOCTOCTG TICTGACCAC ATGACTTTAA AGATATGGAG TATGAAACAA
1251
1301
      CACAATTCTC TCCATCATTT CCACCAACAT AATAAAGAAA TTTATACTAT
1351
      CAAATGGAGT CCAACAGGAC CAGGGACTAA TAATCCAAAT GCCAACCTTA
1401
      TOTAL CAMP TO MAIN SHIP CANAGE PARTS THAT SHIP GEAT CHACAC
      CGAGGGATAT GCATCCATAC CTTGACAAAA CACCAAGAGC CTGTGTACAG
1451
1501
      TGTAGCTITC AGTCCTGATG GCAGGTATCT CCCAAGTGTT TCTTTTCACA
     ANTENDED CANONICS AS ASSCAGACAG GRECTCTAGT TCACAGCTAT
1551
1601
     AGGGGAACAG GYGGAATATT TGAAGTTTGC TGGAATGCAG CAGGAGACAA
1651
     AGTTGGAGCC AGTGCATCAG ATGGTTCAGT TTGTGTATTA GACCTTCGGA
1701
     AATAGegeta etagttggaa gecatggace gactatgaat gtgtacatag
1751
      ccaaaatgag tgtccctgac ccatgtaatg ttatagtccc acttgaacca
1801
      togecagtee aatacageea aatetaaaag aaatatatae atacagtgta
1851
      tataaacaaa attacaccct gaagatgaca gagttttgtc acagcttgtg
1901
      aattetotte accaagtott ggaatetaat etgetotgee eetaaaatag
```

1951 catttagaaq ttttqqatat gaaaaacaga aqaqagaaaa tatacattat 2001 aaaaqcaqta catacatqta ccaqtttttq qatactaaat qacaqccttq 2051 tttctcccct ttgaatcagc agacaccatg gattatattc ttttttccc 2101 ttcagtagtg agcagtttgt atgtacagag aaaatggact tacaaaaact 2151 tgcagcagta gtttgttctt gctttaaaat ttcgtttttg gtttagatta 2201 togatocato aagtaagoga otgaatcagt ttottotta tattttttc 2251 accttttaaa caaaaaattc tttaaaatat tttaatgcat tcttttgaag aggtagatgt ttggtacatt ttatggctcc cagagcatat attcagttgg 2301 tqcatqttqt qqaaqqqqqa attqqaaatt aaatqqaaaa cctatqactt 2351 2401 togtectote aatetotaag acacateagt aaaaaggtat tatgetetgt togttttatt titttatttt gettttttt tittttattt 2451 2501 tootoatoto gettaaatoe aatagtttet tttttgggae atatttetge 2551 caattaaaqa ctagaaqggc acaacttttt ttttaattac catagagaag 2601 atacattaaa aaaaatette tgatgttttg tagccataac taaattatgg taaaaatqtq cactattqtq aaaaqqaqca acqtaqtttt ggqttttttg 2651 ttgtttgttt gttttgcttt gttttttaag agattaaaat gtttctggat 2701 aaggattage ttetegaagt gteeateatt etgtgtanaa gettaaatat 2751 2801 gtaatgtaac caaactccag tattaaaaat ctctcatgtt gttttcttta tacaaagcaa qataacggca tataacactg ccattacatg gcaaaatgtt 2851 tgetacetta gtttaaaaaa caateteaaa caaaagaett getteaaggt 2901 2951 gtttttaaat agcagtgatt cagaattttt tttaatgaaa gtataattgc 3001 actaacette tteetgetge tetgattetg catttgtggt acttgtgact 3051 acotttttc aaatatagat agatttaagc toctaatttt tttttttta gtaatcacta ctatatcatg tettttacte tgtttataat atcaagtatt 3101 ttottaaaga tatagatatt aaacettoto etcatocaac ttagagtaac 3151 3201 atatacagac aaatgattgc atgaggccat gtttatatgt gtgactaata aggottotca toattaacat aatocaggta totcatttot gaagagaata 3251 3301 gtcatcaaat ttatatotog aagattttaa ttaagggaat tgottattgt ttgagettag caaattaata acactattte tgteactaat tattttgagg 3351 3401 ccttttagta ctaaaatttt aacctgtgtt ctaagtagaa actgatttaa cccaagtaat gcagctttga ttgatttcag cattcgttgc tttgctattt 3451 ttacaaaaca gcattgattg aagcaagttt tggttttact aaggtagggt 3501 agcatttgct attggtaaag agaataaata cacttaattt cacaatacat 3551 tottatatot accccaotto ttottagtog ggactatgat actgtaataa 3601 tatttttaaa aatttacatc aagagaggca gtcattcacg atggttttgt 3651 3701 occapetett tttagggttt tggateacat tagagatatt tagaacatat taccetgtga ettacgtagg aaacetaata tgetgagtat etggcaettg 3751 aattootgot tttattgotg gaggtooaca tgtgtggttg acctotgtta 3801 3851 ttotttaaaa aaaaaaaaaa aaaaaaaaaa aaaaa

200000000000000			
161 bp	1544 bp	2180 bp	C 2 1
			20.1:00
	Pe		

FIG.6

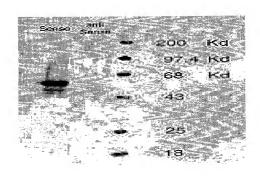


FIG. 7

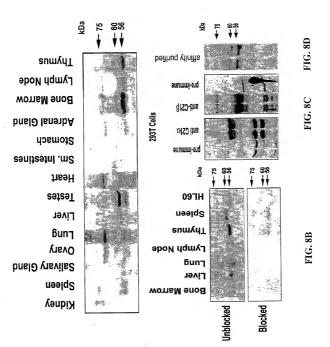
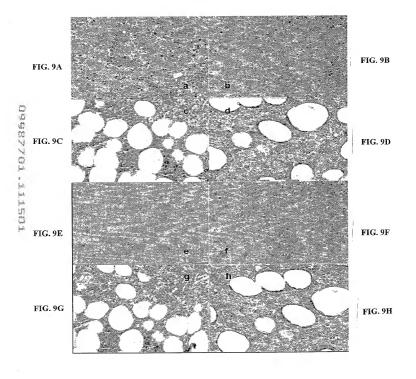


FIG. 8A



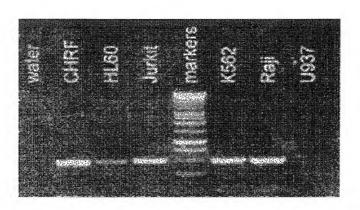


FIG. 10

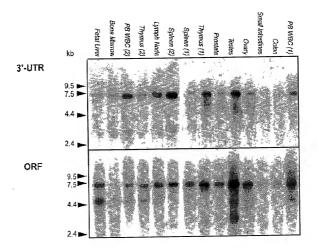
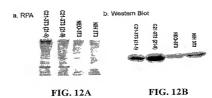
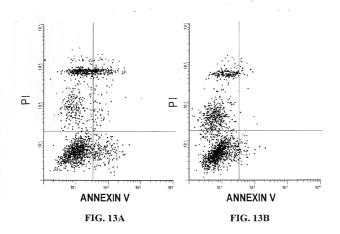


FIG. 11





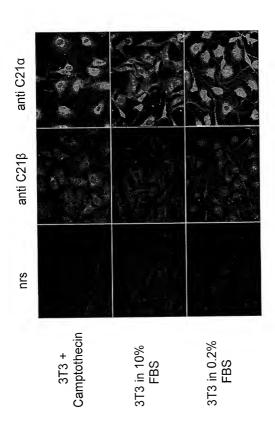


FIG. 14

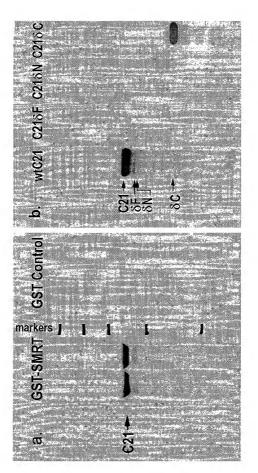


FIG. 15A

FIG. 15B

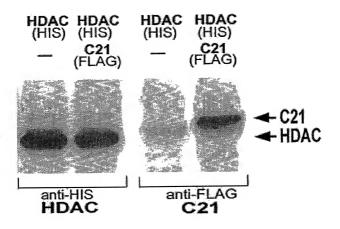


FIG. 16

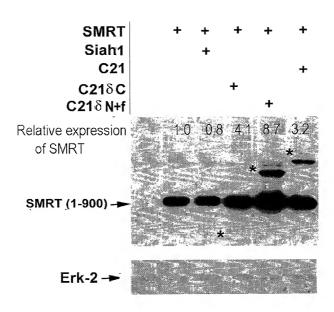


FIG. 17



FIG. 18

Control MG 132

